## SUPPLEMENTAL MATERIAL

14

42

44

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## **Supplemental 1. PROSPERO registration.**

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#### **PROSPERO**

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## Animal review

#### 1. \* Review title.

Give the working title of the review. This must be in English. The title should have the interventions or exposures being reviewed and the associated health or social problems.

Safety and efficacy of cell-based/derived therapies in congenital heart disease; a systematic review and meta analysis of pre-clinical and clinical studies

## 2. Original language title.

For reviews in languages other than English, this field should be used to enter the title in the language of the review. This will be displayed together with the English language title.

#### English

## 3. \* Anticipated or actual start date.

Give the date when the systematic review commenced, or is expected to commence.

## 27/08/2019

## 4. \* Anticipated completion date.

Give the date by which the review is expected to be completed.

### 01/01/2020

## 5. \* Stage of review at time of this submission.

Indicate the stage of progress of the review by ticking the relevant Started and Completed boxes. Additional information may be added in the free text box provided.

Please note: Reviews that have progressed beyond the point of completing data extraction at the time of initial registration are not eligible for inclusion in PROSPERO. Should evidence of incorrect status and/or completion date being supplied at the time of submission come to light, the content of the PROSPERO record will be removed leaving only the title and named contact details and a statement that inaccuracies in the stage of the review date had been identified.

This field should be updated when any amendments are made to a published record and on completion and publication of the review.

The review has not yet started: No

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NHS
National Institute for
Health Research

Review stage	Started	Completed
Preliminary searches	No	Yes
Piloting of the study selection process	No	Yes
Formal screening of search results against eligibility criteria	No	Yes
Data extraction	No	Yes
Risk of bias (quality) assessment	Yes	No
Data analysis	Yes	No

Provide any other relevant information about the stage of the review here (e.g. Funded proposal, protocol not yet finalised).

#### 6. \* Named contact.

The named contact acts as the guarantor for the accuracy of the information presented in the register record.

Alvaro Moreira

## Email salutation (e.g. "Dr Smith" or "Joanne") for correspondence:

Dr Moreira

## 7. \* Named contact email.

Enter the electronic mail address of the named contact.

MoreiraA@uthscsa.edu

## 8. \* Named contact address.

Enter the full postal address for the named contact.

UT Health San AntonioDepartment of Pediatrics, Division of Neonatology7703 Floyd Curl Drive MC 7812San Antonio, TX, USA 78229

# 9. Named contact phone number

Enter the telephone number for the named contact, including international dialling code.

210-567-5226

## 10. \* Organisational affiliation of the review.

Full title of the organisational affiliations for this review and website address if available. This field may be completed as 'none' if the review is not affiliated to any organisation.

UT Health San Antonio

## Organisation web address:

## 11. \* Review team members and their organisational affiliations.

Give the personal details and the organisational affiliations of each member of the review team. Affiliation refers to groups or organisations to which review team members belong. **NOTE: email and country are** 

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now mandatory fields for each person.

Dr John Martinez. UT Health Pediatrics Dr Sarah Zoretic. UT Health Pediatrics

Dr Alvaro Moreira. UT Health San Antonio, Department of Pediatrics, Division of Neonatology

#### 12. \* Funding sources/sponsors.

Give details of the individuals, organisations, groups or other legal entities who take responsibility for initiating, managing, sponsoring and/or financing the review. Any unique identification numbers assigned to the review by the individuals or bodies listed should be included.

Parker B. Francis

## Grant number(s)

#### 13. \* Conflicts of interest.

List any conditions that could lead to actual or perceived undue influence on judgements concerning the main topic investigated in the review.

None

#### 14. Collaborators.

Give the name, affiliation and role of any individuals or organisations who are working on the review but who are not listed as review team members.

#### 15. \* Review question.

Give details of the question to be addressed by the review, clearly and precisely.

Are cell-based/derived therapies both safe and efficacious in clinical trials involving congenital heart disease?

#### Context and rationale

Provide a brief description of the context and rationale of the review, including information on the relevance of your review for human health (max 250 words).

Preclinical studies have established that regenerative therapies show promise as primary/adjunctive therapies for congenital heart disease (CHD). Animal models have demonstrated that regenerative cells are safe and effective. As these therapies have now translated to clinical trials in pediatric CHD, it is imperative to summarize the current findings and identify knowledge gaps that still remain in order optimize translational success. Therefore, the purpose of this systematic review and meta-analysis is twofold: (i) assess the safety, and (ii) efficacy of cell-based/derived therapies in animal models of congenital heart disease.

#### 16. \* Searches.

Give details of the sources to be searched, and any restrictions (e.g. language or publication period). The full search strategy is not required, but may be supplied as a link or attachment.

MEDLINE via PubMed, Scopus, ScienceDirect, Web of Science, Reference lists of included studies, Reference lists of relevant reviews Search dates: no restriction on timeline of search results (initial year-08/26/19)Restrictions on language: no restrictions Publication: no publication date restrictions Will

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searches be re-run prior to final analysis? : yes Will unpublished studies be sought? : no

#### 17. URL to search strategy.

Give a link to the search strategy or an example of a search strategy for a specific database if available (including the keywords that will be used in the search strategies).

Alternatively, upload your search strategy to CRD in pdf format. Please note that by doing so you are consenting to the file being made publicly accessible.

Do not make this file publicly available until the review is complete

#### 18. \* Human disease modelled.

Give a short description of the disease, condition or healthcare domain being modelled.

Congenital heart disease

#### 19. \* Animals/population.

Give summary criteria for the animals being studied by the review, e.g. species, sex, details of disease model. Please include details of both inclusion and exclusion criteria.

#### Inclusion criteria:

Human: Children (newborn-18 years) with congenital heart disease to include adults with history of congenital heart disease receiving cell-based/derived therapies

Animal models of congenital heart disease

#### Exclusion criteria:

Human: Children without congenital heart disease, Adults without congenital heart disease receiving cell-based/derived therapies, Adult models of heart disease

Animal: Animal models without congenital heart disease

## 20. \* Intervention(s), exposure(s).

Give full and clear descriptions of the nature of the interventions or the exposures to be reviewed (e.g. dosage, timing, frequency). Please include details of both inclusion and exclusion criteria.

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NHS National Institute for Health Research

Inclusion criteria:

The following will be used for both human and animal studies. Regenerative cell-based/derived therapy used to treat congenital heart disease. Regenerative cell therapies will be defined as: mesenchymal, embryonic, multipotent, inducible pluripotent cells, progenitor, hematopoietic, umbilical cord, cord blood, c-kit+, secretome, exosome, microRNA, microvesicles, extracellular vesicles.

Exclusion criteria:

The following will be used for both human and animal studies. Non cell-based/derived therapies used to treat congenital heart disease

21. \* Comparator(s)/control.

Where relevant, give details of the type(s) of control interventions against which the experimental condition(s) will be compared (e.g. another intervention or a non-exposed control group). Please include details of both inclusion and exclusion criteria.

Inclusion criteria:

Human: Placebo. Children with congenital heart disease who did not receive cell based therapies.

Animal: Animals in experimental models not subject to cell-based/derived therapies for the treatment of congenital heart disease (placebo and sham).

Exclusion criteria:

Human: Children without congenital heart diseases

Animal: Animals not modeling congenital heart disease

22. \* Study designs to be included.

Give details of the study designs eligible for inclusion in the review. If there are no restrictions on the types of study design eligible for inclusion, or certain study types are excluded, this should be stated. Please include details of both inclusion and exclusion criteria.

Inclusion criteria:

Clinical trials, cohort, case reports

Exclusion criteria:

Articles not assessing outcomes of interest

23. Other selection criteria or limitations applied.

Give details of any other inclusion and exclusion criteria, e.g. publication types (reviews, conference abstracts), publication date, or language restrictions.

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Review articles, book chapters, abstracts will be excluded. No restrictions placed based on publication date or language.

#### 24. \* Outcome measure(s).

Give detail of the outcome measures to be considered for inclusion in the review. Please include details of both inclusion and exclusion criteria.

#### Inclusion criteria:

The following outcome measures will be used for both human and animal studies

Cardiac function, as measured by:

-Right/Left ejection fraction-End diastolic volume-End systolic volume-Tricuspid annular plane systolic excursion-Fractional area change-Fractional shortening Safety: -Mortality -Adverse events with administration (fever, rash, infection, hemodynamic instability, arrhythmia, etc)

#### Exclusion criteria:

Animal or human studies not assessing safety or efficacy (as defined above) after cell-based/derived.

#### 25. N/A.

This question does not apply to systematic reviews of animal studies for human health submissions.

26. \* Study selection and data extraction.

#### Procedure for study selection

Give the procedure for selecting studies for the review, including the screening phases (title and/or title-abstract and/or full-text), the number of researchers involved, and how discrepancies will be resolved. Study selection: a) Two investigators (J. Martinez & S. Zoretic) will independently screen all the abstracts/full texts for the inclusion criteria. b) Differences of opinion in either phase that cannot be resolved by discussion will be resolved by consulting a third investigator (A. Moreira).

## Prioritise the exclusion criteria

Multiple exclusion criteria may apply to an abstract/paper, which can cause discrepancies between reviewers in the reason for exclusion recorded. To avoid this, it is helpful to prioritize the exclusion criteria (e.g. 1) not an animal study; 2) not a myocardial infarction model, etc.) and record the highest ranking applicable criterion as the reason for exclusion. Please sort the exclusion criteria defined in questions 19 to 24. If applicable, do so for each screening phase.

1) in-vitro studies2) studies not including cell-based/derived therapies 3) Human studies not including congenital heart disease models 4) Animal models without congenital heart disease5) Animal or human studies not assessing safety or efficacy (as defined above) after cell-based/derived. 6) Adults without congenital heart disease receiving cell- based/derived therapies 7) Adult models of heart disease 8) Review articles, book chapters, abstracts

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#### Methods for data extraction

Describe methods for data extraction, including the number of reviewers performing data extraction, extraction of data from text and/or graphs, whether and how authors of eligible studies will be contacted to provide missing or additional data, etc.

Data extraction: Study design, methodology, patient demographics, clinical diagnoses, cell characteristics (source, dose, frequency and delivery), cardiac imaging parameters, laboratory values, publication details (author, year, funding, etc), follow up data a) Two investigators (J. Martinez & S. Zoretic) will independently screen all the abstracts/full texts for the inclusion criteria. b) Differences of opinion in either phase that cannot be resolved by discussion will be resolved by consulting a third investigator (A. Moreira). Data will be extracted from text, tables and figures (webplot digitizer). For missing data, will contact authors. Data will be recorded via excel spreadsheet.

## Data to be extracted: study design

Specify the data to be extracted related to characteristics of the study design, e.g. controlled versus crossover, number of experimental groups, etc.

Humans: Number of children in experimental +/- control group, number of experimental groups, phase of clinical trial, cell-based/derived therapies parameters (dose, frequency, route, etc), cardiac assessments (echo, MRI, CT, biomarkers), time points for data collection

Animals: Number of animals in experimental +/- control group, number of experimental groups, cell-based/derived therapies parameters (dose, frequency, route, etc), cardiac assessments (echo, MRI, CT, biomarkers), time points for data collection

## Data to be extracted: animal model

Specify the data to be extracted related to characteristics of the animal model, e.g. species, sex of the animals, etc.

Number of animals in experimental and control groups, power calculation reported, method(s) to induce congenital heart disease, animal species/strain, age, gender, weight and immune status.

### Data to be extracted: intervention of interest

Specify the data to be extracted related to characteristics of the intervention of interest, e.g. dose, timing, etc. Cell type, tissue source, dose, mode of delivery, frequency, timing, passage number

#### Data to be extracted: primary outcome(s)

Define the primary outcome measure(s). For each outcome measure, specify in which format data will be extracted, including the eligible units of measurement, and data type (continuous/dichotomous). A description of any other manipulation or transformation of the extracted data that is planned may be included.

The following outcomes will be assessed in both animal and human studies

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Safety:

-Mortality (dichotomous) -Adverse events with administration (fever, rash, infection, hemodynamic instability, arrhythmia, etc) (dichotomous)

Assessment of cardiac function as measured by:

-Right/Left ejection fraction (%, continuous)-End diastolic volume (mL, continuous)-End systolic volume (mL, continuous) -Tricuspid annular plane systolic excursion (mm, cm, continuous) -Fractional area change (%, continuous) -Fractional shortening (%, continuous)

## Data to be extracted: secondary outcome(s)

Define the secondary outcome measure(s). For each outcome measure, specify in which format data will be extracted, including the eligible units of measurement, and data type (continuous/dichotomous). A description of any other manipulation or transformation of the extracted data that is planned may be included.

n/a

#### Data to be extracted: other

Specify any other data or study characteristics to be extracted, e.g. bibliographical details, such as author, year and language.

Author, year, funding, title, language, contact author email, journal

## 27. \* Risk of bias and/or quality assessment.

State whether and how risk of bias and/or study quality will be assessed. Assessment tools specific for preclinical animal studies include SYRCLE's risk of bias tool and the CAMARADES checklist for study quality

No risk of bias and/or quality assessment planned

No

By use of SYRCLE's risk of bias tool

Yes

By use of SYRCLE's risk of bias tool adapted as follows:

No

By use of the CAMARADES checklist for study quality

No

By use of the CAMARADES checklist for study quality, adapted as follows:

No

Other criteria, namely

Yes

Animal: SYRCLE Risk of bias
Human non randomized: Robins-I

Human randomized: Cochrane Risk of bias

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### Method for risk of bias and/or quality assessment

Give the procedure for the risk of bias and/or quality assessment, including the number of reviewers involved, their contribution, and how discrepancies will be resolved.

Two separate reviewers will assess risk of bias for each study. Discrepancies will be resolved by senior author.

## 28. \* Strategy for data synthesis.

## Planned approach

For each outcome measure, specify whether a quantitative or narrative synthesis is planned and how this decision will be made.

Quantitative synthesis will be preferred method for reporting information, however if 4 studies are assessing a particular outcome we will conduct a narrative explanation as, too few studies will be available to conduct meta-analysis.

If a meta-analysis is planned, please specify the following:

#### Effect measure

For each outcome measure, specify the effect measure to be used (e.g. mean difference, odds ratio etc.).

Animal studies: standardized mean difference

Human studies: odds ratio

## Effect models

For each outcome measure, specify the statistical model of analysis (e.g. random-effects or fixed-effect model).

Random-effects model

## Heterogeneity

Specify the statistical methods to assess heterogeneity (e.g. I², Q). For further guidance please refer to the introduction and practical guide to pre-clinical meta-analysis.

|2

#### Other

Specify other details of the meta-analysis methodology (e.g. correction for multiple testing, correction for multiple use of control group).

n/a

## 29. \* Analysis of subgroups or subsets.

#### Subgroup analyses

Give any planned exploration of subgroups or subsets within the review. 'None planned' is a valid response

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if no subgroup analyses are planned.

Study design: experimental and control groups, congenital heart disease model, measures of safety

measures of function, outcome time

Animal models: species, strain, age, gender

Cell-based/derived therapy source: dose, delivery, timing, frequency, transplant method (allogeneic,

xenogeneic, autologous)

#### Sensitivity

For each outcome measure, specify any sensitivity analyses you propose to perform.

If high heterogeneity is observed (70%), subgroup analyses will be conducted

#### Publication bias

Specify whether an assessment of publication bias is planned. If applicable, specify the method for assessment of publication bias.

funnel plot assessment, Egger's regression

## 30. \* Review type.

#### Type of review

Animal model review

No

Experimental animal exposure review

No

Pre-clinical animal intervention review

Yes

## 31. Language.

Select each country individually to add it to the list below, use the bin icon to remove any added in error. English

There is not an English language summary

## 32. \* Country.

Select the country in which the review is being carried out from the drop down list. For multi-national collaborations select all the countries involved.

United States of America

## 33. Other registration details.

List other places where the systematic review protocol is registered. The name of the organisation and any unique identification number assigned to the review by that organisation should be included.

n/a

## 34. Reference and/or URL for published protocol.

Give the citation and link for the published protocol, if there is one.

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n/a

Give the link to the published protocol.

Alternatively, upload your published protocol to CRD in pdf format. Please note that by doing so you are consenting to the file being made publicly accessible.

#### No I do not make this file publicly available until the review is complete

Please note that the information required in the PROSPERO registration form must be completed in full even if access to a protocol is given.

#### 35. Dissemination plans.

Give brief details of plans for communicating essential messages from the review to the appropriate audiences.

The manuscript will be submitted to a loading journal in field. In addition, a report will be submitted to the funder (Parker B Francis foundation).

## Do you intend to publish the review on completion?

No

## 36. \* Keywords.

Give words or phrases that best describe the review. Separate keywords with a semicolon or new line.

Regenerative medicine, cell-based/derived therapies, stem cells, congenital heart disease, human, clinical trials, animal studies

#### 37. Details of any existing review of the same topic by the same authors.

Give details of earlier versions of the systematic review if an update of an existing review is being registered, including full bibliographic reference if possible.

Previous manuscript focusing on cell-based/derived therapies as a treatment for right ventricular dysfunction is currently being considered for publication.

#### 38. \* Current review status.

Review status should be updated when the review is completed and when it is published. Please provide anticipated publication date

Review\_Ongoing

## 39. Any additional information.

Provide any further information the review team consider relevant to the registration of the review.

## 40. Details of final report/publication(s).

This field should be left empty until details of the completed review are available. Give the full citation for the final report or publication of the systematic review.

Give the link to the published review.

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# Supplemental 2. SYRCLE criteria for animal intervention studies.



## Systematic Review Protocol for Animal Intervention Studies

## FORMAT BY SYRCLE (<u>www.syrcle.nl</u>) Version 2.0 (December 2014)

Item #	Section/Subsection/Item	Description	Check for approval
	A. General		
1.	Title of the review	Safety and efficacy of cell-based therapies in congenital heart disease; a systematic review and meta analysis of pre-clinical and clinical studies	
2.	Authors (names, affiliations, contributions)	John Martinez, MD: conception, study design, search, data collection, protocol writing, manuscript writing Sarah Zoretic, DO: conception, study design, search, data collection, protocol writing, manuscript writing Alvaro Moreira MD, MSc: conception, study design, data collection and analysis, manuscript revision, supervision University of Texas Health San Antonio Department of Pediatrics, Division of Neonatology,	
3.	Other contributors (names, affiliations, contributions)	None	
4.	Contact person + e-mail address	address Alvaro Moreira: MoreiraA@uthscsa.edu	
5.	Funding sources/sponsors	Parker B Francis Foundation	
6.	Conflicts of interest	None	
7.	Date and location of protocol registration	CAMARADES	
8.	Registration number (if applicable)	N/A	
9.	Stage of review at time of registration	Preliminary searches	
	B. Objectives		
	Background		
10.	What is already known about this disease/model/intervention? Why is it important to do this review?	Preclinical studies have established that regenerative therapies show promise as primary/adjunctive therapies for congenital heart disease (CHD). Animal models have demonstrated that regenerative cells are safe and effective. As these therapies have now translated to clinical trials in pediatric CHD, it is imperative to summarize the current findings and identify knowledge	

		gaps that still remain in order optimize translational success. Therefore, the purpose of this systematic review and meta-analysis is twofold: (i) assess the safety, and (ii) efficacy of cell-based/derived therapies in animal models of congenital heart disease.			
	Research question				
11.	Specify the disease/health problem of interest	Congenital heart disease: Hypoplastic left heart syndrome, Tricuspid atresia, Single ventricle physiology, Transposition of great arteries, Tetralogy of Fallot, Pulmonary Atresia, Anomalous pulmonary venous return, Double outlet right ventricle, Single inlet ventricle, Coarctation of aorta, Interrupted aortic arch, Ebstein's anomaly.			
12.	Specify the population/species studied	Animal models of congenital heart disease (as listed above)			
13.	Specify the intervention/exposure	Cell-based/derived therapies: mesenchymal, embryonic, multipotent, inducible pluripotent cells, progenitor, hematopoietic, umbilical cord, cord blood, c-kit+, secretome, exosome, microRNA, microvesicle, extracellular vesicle.			
14.	Specify the control population				
15.	Specify the outcome measures	Primary Outcome: Safety and cardiac function (refer to number 26)			
16.	State your research question (based on items 11-15)	Are cell-based/derived therapies both safe and efficacious in experimental models of congenital heart disease?			
	C. Methods				
	Search and study identification	V			
17.	Identify literature databases to search (e.g. Pubmed, Embase, Web of science)	X MEDLINE via PubMed XWeb of Science  XSCOPUS □EMBASE  XOther, namely: Science direct □Specific journal(s), namely:			
18.	Define electronic search strategies (e.g. use the step by step search guide <sup>15</sup> and animal search filters <sup>20, 21</sup> )	When available, please add a supplementary file containing your search strategy: [insert file name]			
	guide and animal search filters——)				
19.	Identify other sources for study identification	XReference lists of included studies ☐Books XReference lists of relevant reviews ☐Conference proceedings, namely: ☐Contacting authors/ organisations, namely: ☐Other, namely:			
19.	Identify other sources for study	XReference lists of included studies ☐Books XReference lists of relevant reviews ☐Conference proceedings, namely: ☐Contacting authors/ organisations, namely:			

Define screening phases (e.g. prescreening based on title/abstract, full text screening based on title/abstract, full text screening, both)  Specify (a) the number of reviewers per screening phase and (b) how discrepancies will be resolved phase and (b) how discrepancies will be resolved phase and exclusion criteria based on:  Type of study (design)  First phase: screening by title and abstract Second phase: full text screening of eligible articles Full text studies that do not meet inclusion will be incorporated into the flow diagram with reasons for exclusion a) Two investigators (J. Martinez & S. Zoretic) will independently screen all the abstracts/full texts for the inclusion criteria. b) Differences of opinion in either phase that cannot be resolved by discussion will be resolved by consulting a third investigator (A. Moreira).  Define all inclusion and exclusion criteria based on:  Inclusion criteria: pre-clinical studies Exclusion criteria: non-intervention studies, no control group, co-intervention studies	
21. screening based on title/abstract, full text studies that do not meet inclusion will be incorporated into the flow diagram with reasons for exclusion  a) Two investigators (J. Martinez & S. Zoretic) will independently screen all the abstracts/full texts for the inclusion criteria. b) Differences of opinion in either phase that cannot be resolved by discussion will be resolved by consulting a third investigator (A. Moreira).  Define all inclusion and exclusion criteria based on:  Inclusion criteria: pre-clinical studies Exclusion criteria: non-intervention studies, no control	
text screening, both)  incorporated into the flow diagram with reasons for exclusion  a) Two investigators (J. Martinez & S. Zoretic) will independently screen all the abstracts/full texts for the inclusion criteria. b) Differences of opinion in either phase that cannot be resolved by discussion will be resolved by consulting a third investigator (A. Moreira).  Define all inclusion and exclusion criteria based on:  Inclusion criteria: pre-clinical studies Exclusion criteria: non-intervention studies, no control	
exclusion  a) Two investigators (J. Martinez & S. Zoretic) will independently screen all the abstracts/full texts for the inclusion criteria. b) Differences of opinion in either phase that cannot be resolved by discussion will be resolved by consulting a third investigator (A. Moreira).  Define all inclusion and exclusion criteria based on:  Inclusion criteria: pre-clinical studies Exclusion criteria: non-intervention studies, no control	
a) Two investigators (J. Martinez & S. Zoretic) will independently screen all the abstracts/full texts for the inclusion criteria. b) Differences of opinion in either phase that cannot be resolved by discussion will be resolved by consulting a third investigator (A. Moreira).  Define all inclusion and exclusion criteria based on:  Inclusion criteria: pre-clinical studies Exclusion criteria: non-intervention studies, no control	
Specify (a) the number of reviewers per screening phase and (b) how discrepancies will be resolved that cannot be resolved by discussion will be resolved by consulting a third investigator (A. Moreira).  Define all inclusion and exclusion criteria based on:  Inclusion criteria: pre-clinical studies Exclusion criteria: non-intervention studies, no control	
22. per screening phase and (b) how discrepancies will be resolved that cannot be resolved by discussion will be resolved by consulting a third investigator (A. Moreira).  Define all inclusion and exclusion criteria based on:  Inclusion criteria: pre-clinical studies Exclusion criteria: non-intervention studies, no control	
discrepancies will be resolved that cannot be resolved by discussion will be resolved by consulting a third investigator (A. Moreira).  Define all inclusion and exclusion criteria based on:  Inclusion criteria: pre-clinical studies Exclusion criteria: non-intervention studies, no control	
consulting a third investigator (A. Moreira).  Define all inclusion and exclusion criteria based on:  Inclusion criteria: pre-clinical studies Exclusion criteria: non-intervention studies, no control	
Define all inclusion and exclusion criteria based on:  Inclusion criteria: pre-clinical studies  Type of study (design)  Exclusion criteria: non-intervention studies, no control	
23. Type of study (design)  Inclusion criteria: pre-clinical studies Exclusion criteria: non-intervention studies, no control	
23. Type of study (design) Exclusion criteria: non-intervention studies, no control	
group, co-intervention studies	
Inclusion criteria: animal models of congenital heart	
Type of animals/population (e.g. age, disease, all genders	
gender, disease model) Exclusion criteria: humans, in-vitro, non-pediatric models	
of heart disease	
Inclusion criteria: administration of cell-based/derived	
therapy- all dosages, timing, and frequency; cells may be	
Type of intervention (e.g. dosage, derived from any tissue source	
timing, frequency) Exclusion criteria: Cardiac administration of cell-	
based/derived therapy assessing for variables other than	
safety or effect on function.	
Cardiac function as measured by:	
-Right/Left ejection fraction	
-End diastolic volume	
-End systolic volume	
-Tricuspid annular plane systolic excursion	
-Fractional area change	
26. Outcome measures -Fractional shortening	
20. Outcome measures	
Safety:	
-Mortality	
-Adverse events with administration (fever, rash, infection,	
hemodynamic instability, arrhythmias, etc)	
Exclusion criteria:	
27. Language restrictions Inclusion criteria: English and Spanish	
Exclusion criteria: All other languages	
28. Publication date restrictions Inclusion criteria: no publication date restrictions	
Exclusion criteria:	
29. Other	
Exclusion criteria:	
Sort and prioritize your exclusion  Selection phase: title and abstract screening	
30. Criteria per selection phase 1. Not a primary study	
2. Not an in vivo animal study	

	1	T	
		3. Not congenital heart disease	
		4. No cell based/derived therapy use	
		5. Adult animal	
		Selection phase: full text screening	
		1. Not a primary study	
		2. Not an in vivo animal study	
		3. Not congenital heart disease	
		4. No cell based/derived therapy use	
		5. No assessment of safety, effect on ventricular function	
		5. No control group	
		6. Co-intervention studies	
	Study characteristics to be extracted (for	or assessment of external validity, reporting quality)	
31.	Study ID (e.g. authors, year)	Authors, journal, title, year, language, contact author e-mail	
	Study design characteristics (e.g.	Number of animals in experimental and control groups,	
32.	experimental groups, number of	reporting of randomization process, power calculation	
	animals)	reported, method(s) to induce congenital heart disease	
33.	Animal model characteristics (e.g.	Animal species, strain, age, gender, weight, and immune	
33.	species, gender, disease induction)	status	
34.	Intervention characteristics (e.g.	Source, dose, delivery, timing, and frequency of	
34.	intervention, timing, duration)	intervention	
35.	Outcome measures	Assessment of safety as defined through mortality or	
55.	Outcome measures	occurrence of adverse events upon administration.	
36.	Other (e.g. drop-outs)	Assessment of cardiac function through measures as	
30.		noted above	
	Assessment risk of bias (internal validity	y) or study quality	
37.	Specify (a) the number of reviewers assessing the risk of bias/study quality in each study and (b) how discrepancies will be resolved	a) Two investigators (J. Martinez & S. Zoretic) will independently screen all the abstracts/full texts for the inclusion criteria. b) Differences of opinion in either phase that cannot be resolved by discussion will be resolved by consulting a third investigator (A. Moreira)	
	Define criteria to assess (a) the	X By use of SYRCLE's Risk of Bias tool <sup>4</sup>	
	internal validity of included studies	☐ By use of SYRCLE's Risk of Bias tool, adapted as follows:	
38.	(e.g. selection, performance,	☐ By use of CAMARADES' study quality checklist, e.g <sup>22</sup>	
	detection and attrition bias) and/or (b) other study quality measures (e.g. reporting quality, power)	By use of CAMARADES' study quality checklist, adapted as follows:	
		☐ Other criteria, namely:	
	Collection of outcome data		
	For each outcome measure, define	All outcome measures will be expressed through study	
20	the type of data to be extracted (e.g.	units of measure, values expressed as continuous	
39.	continuous/dichotomous, unit of	measures will be recorded as means +/- SD, SEM or	
	measurement)	median +/- IQR	

40.	Methods for data extraction/retrieval (e.g. first extraction from graphs using a digital screen ruler, then contacting authors)	Extraction from text, tables, and figures (GetData graph digitizer 2.26) Contact authors in case of missing data	
41.	Specify (a) the number of reviewers extracting data and (b) how discrepancies will be resolved	a) Two investigators (J. Martinez & S. Zoretic) will independently screen all the abstracts/full texts for the inclusion criteria. b) Differences of opinion in either phase that cannot be resolved by discussion will be resolved by consulting a third investigator (A. Moreira)	
	Data analysis/synthesis		
42.	Specify (per outcome measure) how you are planning to combine/compare the data (e.g. descriptive summary, meta-analysis)	For sufficient data, we will conduct a meta-analysis for eligible studies. If insufficient data to measure outcomes, we will provide a descriptive summary of study results	
43.	Specify (per outcome measure) how it will be decided whether a meta-analysis will be performed	A minimum of 4 articles for the same outcome is required. High heterogeneity is expected between studies due to differences in the study designs. We will perform a meta-regression analysis to investigate sources of heterogeneity.	
	If a meta-analysis seems feasible/sensib	ble, specify (for each outcome measure):	
44.	The effect measure to be used (e.g. mean difference, standardized mean difference, risk ratio, odds ratio)	Continuous outcomes will be analysed using standardized mean differences (95% CI)	
45.	The statistical model of analysis (e.g. random or fixed effects model)	Random effects model	
46.	The statistical methods to assess heterogeneity (e.g. I <sup>2</sup> , Q)	I <sup>2</sup>	
47.	Which study characteristics will be examined as potential source of heterogeneity (subgroup analysis)	Study design: experimental and control groups, congenital heart disease model, measures of safety, measures of cardiac function.  Animal model: species, strain, age, gender Cell based/derived therapy source: dose, delivery, timing, frequency, transplant method (allogeneic, xenogeneic, autologous, etc.)	
48.	Any sensitivity analyses you propose to perform	If high heterogeneity is observed (≥70%), subgroup analyses will be conducted	
49.	Other details meta-analysis (e.g. correction for multiple testing, correction for multiple use of control group)	N/A	
50.	The method for assessment of publication bias	Funnel plot assessment Egger's regression	
John Martinez MD Sarah Zoretic DO Alvaro Moreira MD, MSc  Date:			

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Supplemental 3. Database search terms.

## **Database Search Terms:**

 ("regenerative" OR "stem cell" OR "stromal cell" OR "mesenchymal" OR "embryonic" OR "pluripotent" OR "multipotent" OR "inducible pluripotent" OR "progenitor" OR "hematopoietic" OR "umbilical cord" OR "cord blood" OR "microparticle" OR "extracellular vessicles") AND ("tetralogy of fallot" or "single ventricle" or "transposition of great arteries" or "anomalous pulmonary venous" or "tricuspid atresia" or "truncus arteriosus" or "hypoplastic left heart" or "ebstein" or "double outlet right ventricle" or "hypoplastic right heart" or "pulmonary atresia" or "coarctation of aorta" or "interrupted aortic arch" or "single inlet ventricle")

## Supplemental 4. List of included studies.

104 105 1. Agarwal U, Sr

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   Implantation of Thymus Stem Cell Engineered Graft in Growing Swine. *JACC Basic to* Transl Sci 2019;4:364–384.
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# **Supplemental 5. PRISMA Checklist.**



# PRISMA 2009 Checklist

Section/topic	#	Checklist item	Reported on page #
TITLE			
Title	1	Identify the report as a systematic review, meta-analysis, or both.	2
ABSTRACT			
Structured summary	2	Provide a structured summary including, as applicable: background; objectives; data sources; study eligibility criteria, participants, and interventions; study appraisal and synthesis methods; results; limitations; conclusions and implications of key findings; systematic review registration number.	3
INTRODUCTION			
Rationale	3	Describe the rationale for the review in the context of what is already known.	4
Objectives	4	Provide an explicit statement of questions being addressed with reference to participants, interventions, comparisons, outcomes, and study design (PICOS).	4
METHODS			
Protocol and registration	5	Indicate if a review protocol exists, if and where it can be accessed (e.g., Web address), and, if available, provide registration information including registration number.	5
Eligibility criteria	6 Specify study characteristics (e.g., PICOS, length of follow-up) and report characteristics (e.g., years considered, language, publication status) used as criteria for eligibility, giving rationale.		5-6
nformation sources 7 Describe all information sources (e.g., databases with dates of coverage, contact with study authors to identify additional studies) in the search and date last searched.		5	
Search 8 Present full electronic search strategy for at least one database, including any limits used, such that it could be repeated.		5	
Study selection  9 State the process for selecting studies (i.e., screening, eligibility, included in systematic review, and, if applicable, included in the meta-analysis).		5	
Data collection process 10 Describe method of data extraction from reports (e.g., piloted forms, independently, in duplicate) and any processes for obtaining and confirming data from investigators.		5-6	
Data items	Data items  11 List and define all variables for which data were sought (e.g., PICOS, funding sources) and any assumptions and simplifications made.		5-6
Risk of bias in individual studies	12	Describe methods used for assessing risk of bias of individual studies (including specification of whether this was done at the study or outcome level), and how this information is to be used in any data synthesis.	8
Summary measures	13	State the principal summary measures (e.g., risk ratio, difference in means).	7-8
Synthesis of results	14	Describe the methods of handling data and combining results of studies, if done, including measures of consistency (e.g., I <sub>2</sub> ) for each meta-analysis.	7-8

Page 1 of 2



## PRISMA 2009 Checklist

Section/topic	#	Checklist item	Reported on page #
Risk of bias across studies	15	cify any assessment of risk of bias that may affect the cumulative evidence (e.g., publication bias, selective orting within studies).	
Additional analyses	16	Describe methods of additional analyses (e.g., sensitivity or subgroup analyses, meta-regression), if done, indicating which were pre-specified.	7-8
RESULTS			
Study selection	17	Give numbers of studies screened, assessed for eligibility, and included in the review, with reasons for exclusions at each stage, ideally with a flow diagram.	9
Study characteristics	18	For each study, present characteristics for which data were extracted (e.g., study size, PICOS, follow-up period) and provide the citations.	9, 11
Risk of bias within studies	19	Present data on risk of bias of each study and, if available, any outcome level assessment (see item 12).	13-14
Results of individual studies	Its of individual studies 20 For all outcomes considered (benefits or harms), present, for each study: (a) simple summary data for each intervention group (b) effect estimates and confidence intervals, ideally with a forest plot.		9-13
Synthesis of results	21	Present results of each meta-analysis done, including confidence intervals and measures of consistency.	
Risk of bias across studies	22	Present results of any assessment of risk of bias across studies (see Item 15).	13-14
Additional analysis 23 Give results of additional analyses, if done (e.g., sensitivity or subgroup analyses, meta-regression [see Item 16]).		10-11, 12-13	
DISCUSSION			
Summary of evidence	24	Summarize the main findings including the strength of evidence for each main outcome; consider their relevance to key groups (e.g., healthcare providers, users, and policy makers).	15-18
Limitations	nitations 25 Discuss limitations at study and outcome level (e.g., risk of bias), and at review-level (e.g., incomplete retrieval of identified research, reporting bias).		18
Conclusions	26	Provide a general interpretation of the results in the context of other evidence, and implications for future research.	19
FUNDING	<u>L</u>		
Funding	27	Describe sources of funding for the systematic review and other support (e.g., supply of data); role of funders for the systematic review.	19

From: Moher D, Liberati A, Tetzlaff J, Altman DG, The PRISMA Group (2009). Preferred Reporting Items for Systematic Reviews and Meta-Analyses: The PRISMA Statement. PLoS Med 6(7): e1000097. doi:10.1371/journal.pmed1000097

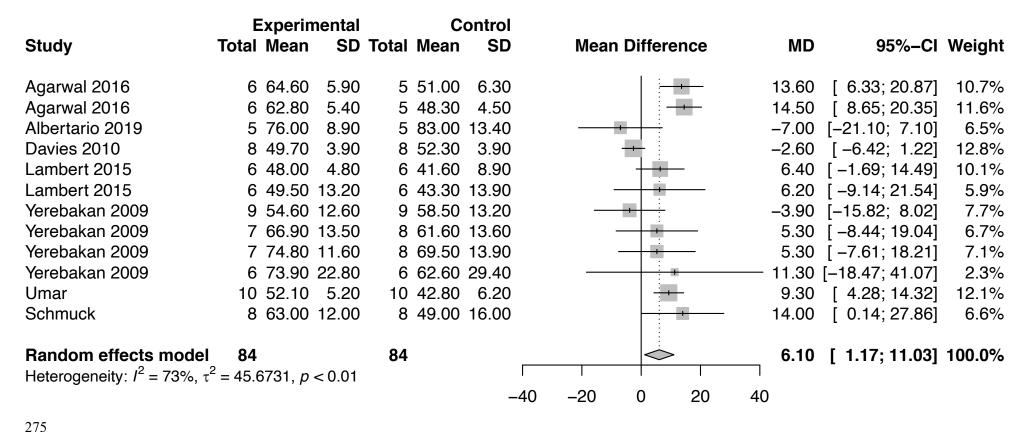
For more information, visit: www.prisma-statement.org.

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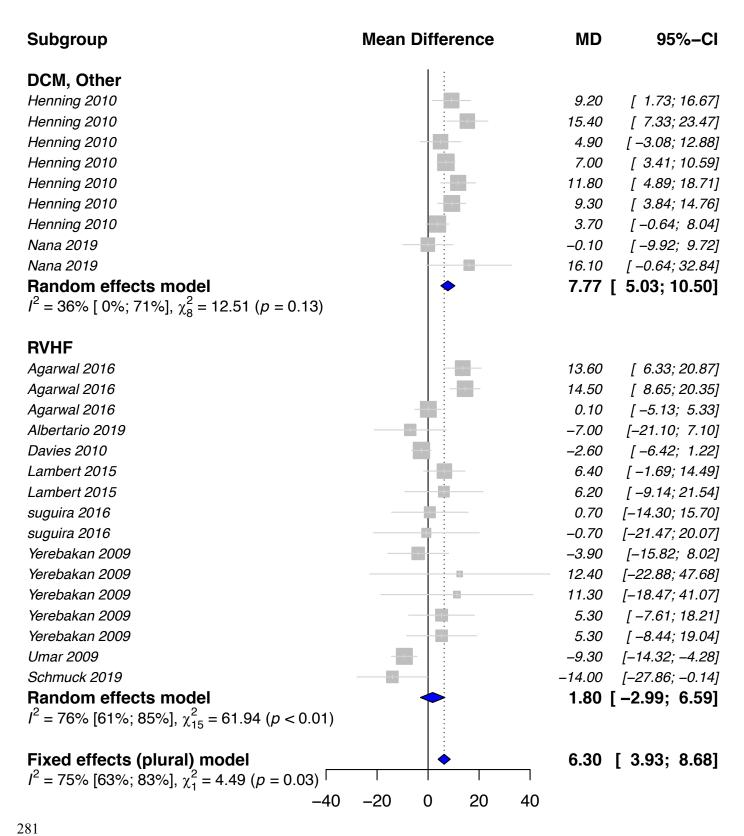
**Supplemental 6. Effect size of regenerative cell on animal ejection fraction.** Forest plots demonstrating MD and 95% CI for A) Left ventricular ejection fraction; cell-based n=172; control=177; p <0.0001. B) Right ventricular ejection fraction; cell-based n=84; control n=84; p=0.02. C) Disease model; cell-based n= 256; control n=261; RVHF, p=0.01; DCM, p<0.0001.

	Experimental	Control		
Study	Total Mean SD	Total Mean SD	Mean Difference	MD 95%-CI Weight
Henning 2010	22 53.50 8.00	23 41.70 14.80	: -	11.80 [ 4.89; 18.71] 9.7%
Henning 2010	22 49.80 16.90	23 44.90 9.10	<del>  = </del>	4.90 [-3.08; 12.88] 7.7%
Henning 2010	12 57.00 9.40	10 41.60 9.80	<del></del>	15.40 [ 7.33; 23.47] 7.6%
Henning 2010	12 52.30 11.80	10 43.10 5.40	- <del>i=</del>	9.20 [ 1.73; 16.67] 8.6%
Henning 2010	22 87.50 6.10	23 83.80 8.60	<del>                                     </del>	3.70 [ -0.64; 8.04] 18.4%
Henning 2010	22 76.00 11.30	23 66.70 6.70	-	9.30 [ 3.84; 14.76] 13.7%
Henning 2010	22 64.20 6.10	23 57.20 6.20	+	7.00 [ 3.41; 10.59] 22.4%
Nana 2019	6 81.50 7.40	8 65.40 22.60		16.10 [ -0.64; 32.84] 2.0%
Nana 2019	6 78.30 7.40	8 78.40 11.30	<del>- + :</del>	-0.10 [-9.92; 9.72] 5.4%
Suguira 2016	10 74.30 21.40	10 73.60 11.30	<del>-  -  -</del>	0.70 [-14.30; 15.70] 2.5%
Suguira 2016	10 72.90 28.20	10 73.60 18.10	<del></del>	-0.70 [-21.47; 20.07] 1.3%
Yerebakan 2009	6 72.70 7.80	6 60.30 43.40		- 12.40 [ <i>-</i> 22.88; 47.68]     0.5%
Random effects mode	l 172	177	<b>♦</b>	7.41 [ 4.96; 9.85] 100.0%
Heterogeneity: $I^2 = 21\%$ ,	$\tau^2 = 3.5835,  p = 0.24$	Г		
		-4	0 -20 0 20 40	

Supplemental 6. A) Cell-based effect on animal LVEF.

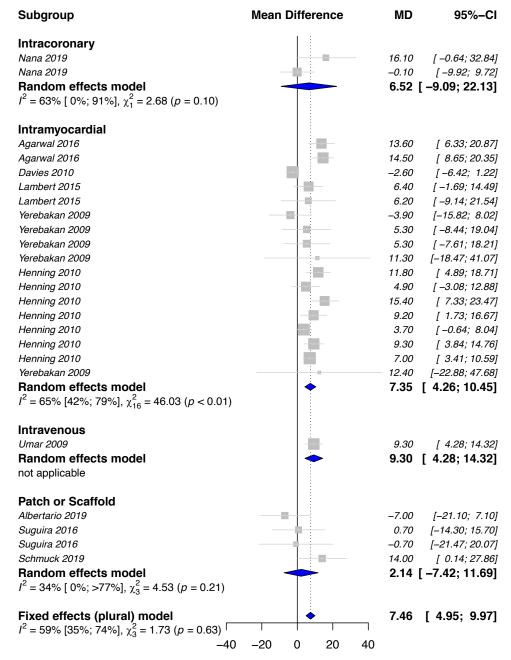


Supplemental 6. B) Cell-based effect on animal RVEF.

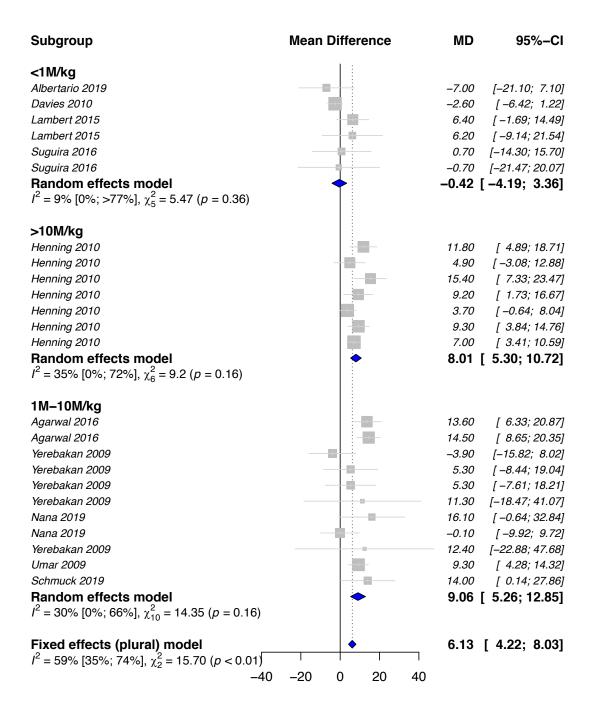


Supplemental 6. C) Cell-based effect on animal ejection fraction by disease model.

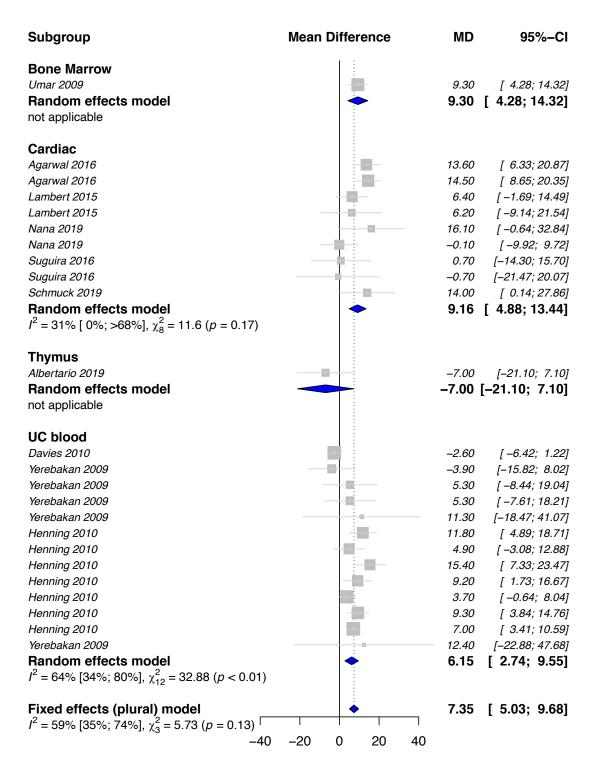
**Supplemental 7. Subgroup analysis of regenerative cell effect size on animal ejection fraction**. Forest plots demonstrating MD and 95% CI for A) Route of delivery, p<0.00001 for intramyocardial injection. B) Dose, p<0.00001 for 1-10 M. C) Tissue Source, p<0.0001 for cardiac; p=0.0003 for bone marrow. D) Timing of delivery, p<0.0001 for 1 week–1 month. E) autologous vs. non-autologous sources, p<0.0001 (non-autologous). Cell-based n=256; Control n=261.



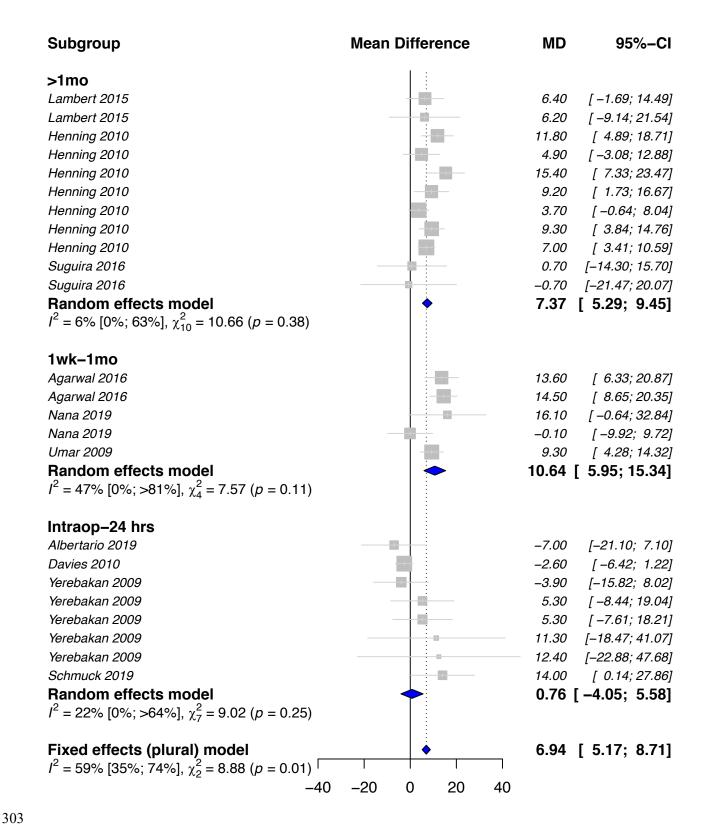
Supplemental 7. A) Cell-based effect on animal ejection fraction by route of delivery.



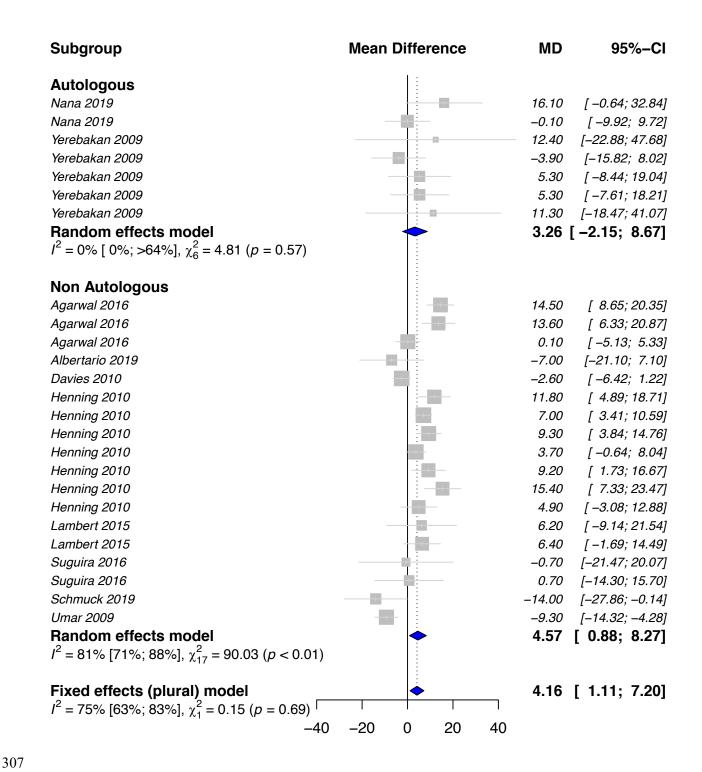
Supplemental 7. B) Cell-based effect on animal ejection fraction by dose.



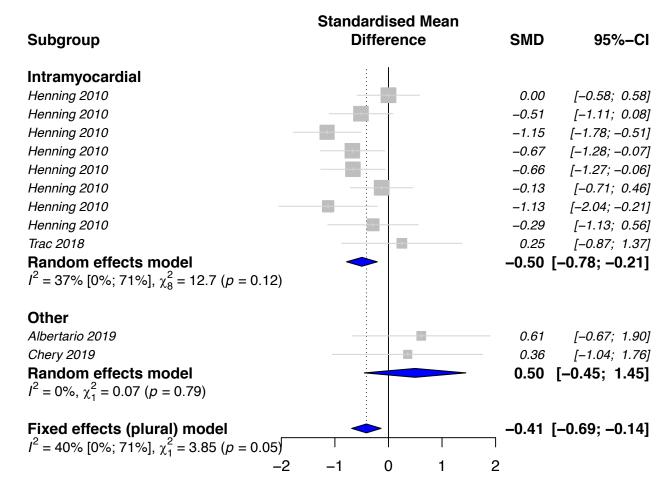
Supplemental 7. C) Cell-based effect on animal ejection fraction by tissue source.



Supplemental 7. D) Cell-based effect on animal ejection fraction by timing of delivery.



Supplemental 7. E) Cell-based effect on animal ejection fraction by autologous vs. non-autologous sources.

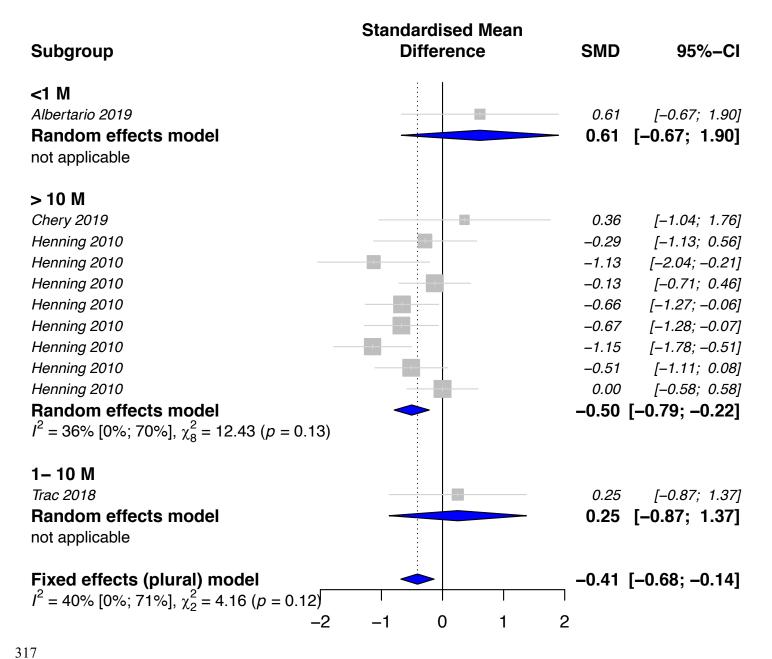


Supplemental 8. A) Cell-based effect on animal FS by route of delivery.

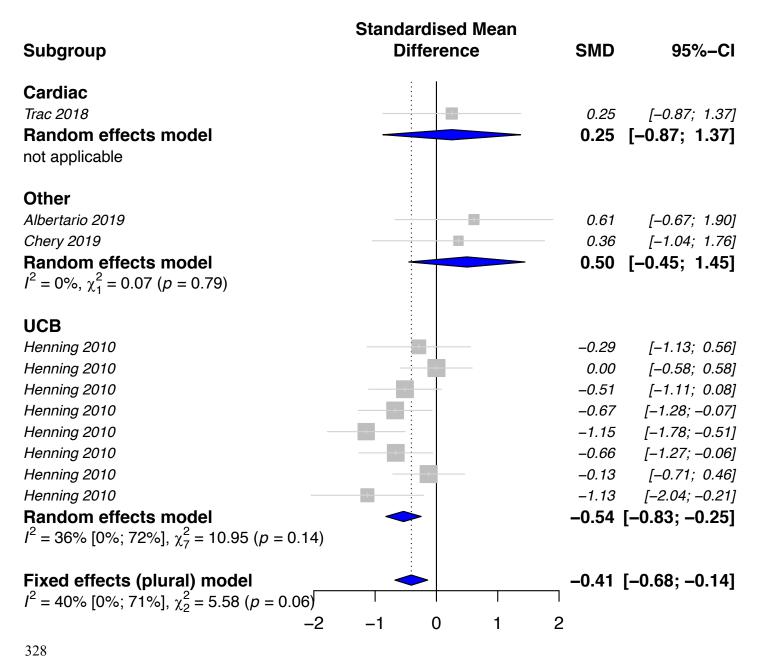
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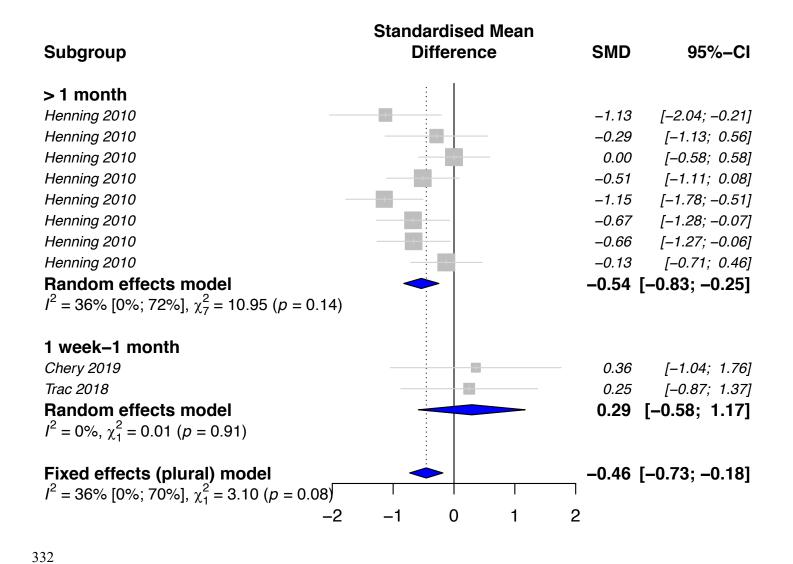
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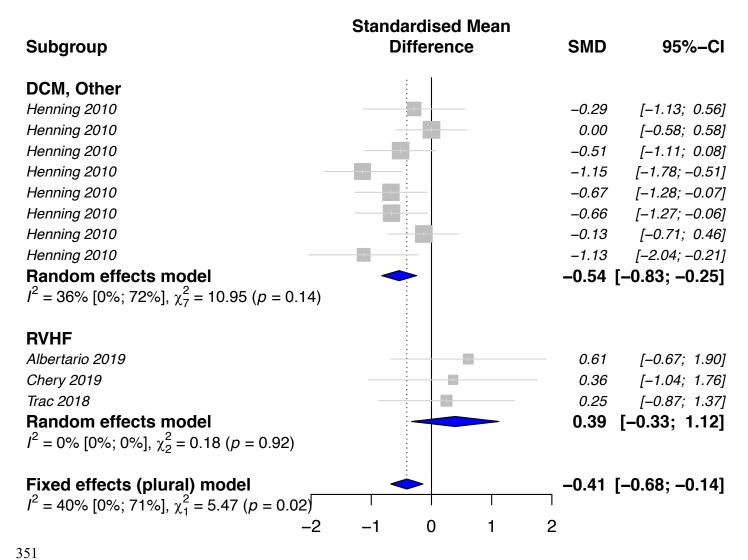
Supplemental 8. B) Cell-based effect on animal FS by dose.



Supplemental 8. C) Cell-based effect on animal FS by tissue source.



Supplemental 8. D) Cell-based effect on animal FS by timing of delivery.



Supplemental 8. E) Cell-based effect on fractional shortening by disease model.

Supplemental 9. Effect size of regenerative cell on additional measures of animal cardiac function. Forest

- plots demonstrating MD and 95% CI for A) Fractional area change, p=0.05; cell-based n= 33; control n=30. B)
- End diastolic volume, p=0.48; cell-based n=67; control n=61. C) End systolic volume, p=0.60; cell-based n=72;
- control n=66. D) Tricuspid annular plane systolic excursion, p=0.55; cell-based n=33; control n=58.

		Expe	rimental			Control				
Study	Total	Mean	SD	Total	Mean	SD	Mean Difference	MD	95%-CI	Weight
Albertario 2019	5	47.10	7.8000	5	49.20	6.5000	<del>-    </del>	-2.10	[-11.00; 6.80]	17.0%
Trach 2018	8	39.80	14.9000	5	27.30	10.5000	<del>                                     </del>	12.50	[ -1.33; 26.33]	13.6%
Wehman 2016	5	42.20	6.5000	5	46.00	2.5000		-3.80	[-9.90; 2.30]	18.7%
Wehman 2016	5	47.80	7.6000	5	29.80	10.7000	-	18.00	[ 6.50; 29.50]	15.2%
Wehman 2017	5	53.40	6.3000	5	46.00	2.5000	<del></del>	7.40	[ 1.46; 13.34]	18.8%
Wehman 2017	5	53.20	1.3000	5	29.90	10.5000	-	23.30	[ 14.03; 32.57]	16.7%
Random effects model Prediction interval	33			30				8.65	[-0.11; 17.42] [-21.38; 38.69]	100.0%
Heterogeneity: $I^2 = 84\%$ , $\tau^2$	<sup>2</sup> – 97 (	1780 n	~ 0 01						[ 21.50, 50.09]	
1 10 to 10 go 110 ity . 1 = 04 /0, t	- 37.0	που, ρ	<b>~</b> 0.01				-30-20-10 0 10 20 30			

Supplemental 9. A) Cell-based effect on animal FAC.

359

	Experimenta	Control	Standardised Mean	
Study	Total Mean SD	Total Mean SD	Difference	SMD 95%-CI Weight
Davies 2010	8 27.90 18.70	8 29.40 19.80		-0.07 [-1.05; 0.91] 12.7%
Lambert 2015	6 166.00 112.70	6 148.00 58.80		- 0.18 [-0.95; 1.32] 9.4%
Lambert 2015	6 192.00 134.70	6 208.00 144.50		-0.11 [-1.24; 1.03] 9.5%
Nana 2019	8 220.00 138.60	6 213.00 193.00		0.04 [-1.02; 1.10] 10.9%
Nana 2019	8 288.00 172.50	6 225.00 154.30		- 0.36 [-0.71; 1.43] 10.6%
Yerebakan 2009	6 21.60 23.00	6 26.60 17.10 -	-	-0.23 [-1.36; 0.91] 9.4%
Yerebakan 2009	9 103.90 32.40	9 90.20 52.80		0.30 [-0.63; 1.23] 14.1%
Yerebakan 2009	8 47.80 20.10	7 49.90 17.70		-0.10 [-1.12; 0.91] 11.8%
Yerebakan 2009	8 27.90 11.00	7 31.10 4.50 -	-	-0.35 [-1.37; 0.68] 11.6%
Random effects mode Heterogeneity: $I^2 = 0\%$ , $\tau^2$	_	61		0.01 [-0.34; 0.36] 100.0%
<b>3 3</b>	, 1		-1 -0.5 0 0.5 1	

Supplemental 9. B) Cell-based effects on animal EDV.

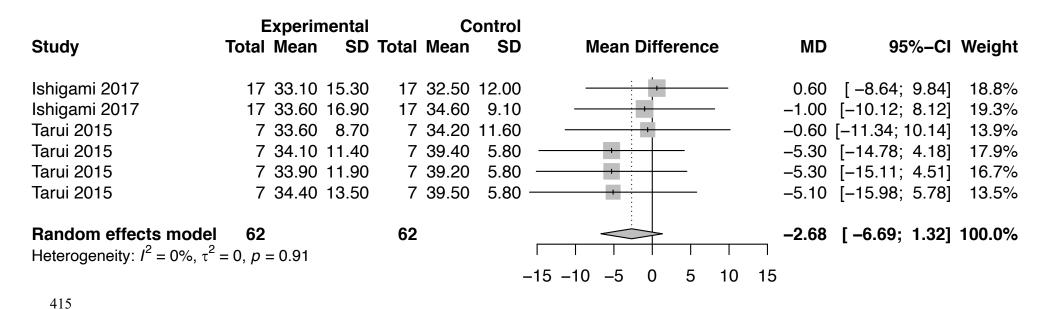
		Experi	mental		C	ontrol	Standardised Mean			
Study	Total	Mean	SD	Total	Mean	SD	Difference	SMD	95%-CI	Weight
Albertario 2019	5	10.00	13.40	5	14.00	2.20		-0.38	[–1.63; 0.88]	7.1%
Davies 2010	8	18.60	10.20	8	17.70	12.70		0.07	[-0.91; 1.05]	11.7%
Lambert 2015	6	67.00	49.00	6	80.00	95.50		-0.16	[-1.29; 0.98]	8.8%
Lambert 2015	6	103.00	88.20	6	83.00	90.60		0.21	[-0.93; 1.34]	8.7%
Nana 2019	8	45.00	39.60	6	47.00	51.40		-0.04	[-1.10; 1.02]	10.0%
Nana 2019	8	100.00	679.00	6	43.00	36.70	<u> </u>	0.10	[-0.96; 1.16]	10.0%
Yerebakan 2009	8	19.90	13.60	7	18.20	10.30		0.13	[-0.88; 1.15]	10.9%
Yerebakan 2009	8	9.50	7.90	7	7.80	3.40		0.26	[-0.76; 1.28]	10.8%
Yerebakan 2009	6	9.00	6.40	6	7.20	9.30		0.21	[-0.93; 1.34]	8.7%
Yerebakan 2009	9	43.20	22.20	9	38.90	27.90		0.16	[-0.76; 1.09]	13.1%
Random effects model		1.00		66				0.07 [	[-0.26; 0.41]	100.0%
Heterogeneity: $I^2 = 0\%$ , $\tau^2$	= 0, $\mu$ =	1.00				_	-1.5 –1 –0.5 0 0.5 1 1.5	5		

Supplemental 9. C) Cell-based effect on animal ESV.

	Ex	perim	ental		Co	ntrol	Standardised Mean			
Study	Total	Mean	SD	Total	Mean	SD	Difference	SMD	95%-CI	Weight
							· I —			
Agarwal 2016	4	1.80	0.20	9	1.70	0.30	<del>-                                    </del>	0.34	[-0.85; 1.52]	13.5%
Agarwal 2016	4	1.60	0.40	9	1.90	0.60		-0.50	[-1.71; 0.70]	13.3%
Trac 2018	5	1.50	0.70	8	0.80	1.40		0.55	[-0.60; 1.69]	14.6%
Trac 2018	5	1.60	1.80	8	1.30	1.10		0.20	[-0.92; 1.32]	15.2%
Trac 2018	5	1.20	1.10	8	1.60	1.10	<del></del>	-0.34	[-1.47; 0.79]	15.0%
Trac 2018	5	1.50	0.50	8	1.80	0.80		-0.40	[-1.53; 0.74]	14.9%
Trac 2018	5	1.50	0.70	8	2.10	0.60	<del> </del>	-0.87	[-2.06; 0.31]	13.5%
							<u> </u>			100.00/
Random effects model				58				0.14	[-0.58; 0.30]	100.0%
Heterogeneity: $I^2 = 0\%$ , $\tau^2$	= 0, p =	= 0.61						I		
						-	-2 –1 0 1	2		

Supplemental 9. D) Cell-based effect on Animal TAPSE.

**Supplemental 10. Effect size of regenerative cell on additional measures of human cardiac function.** Forest plots demonstrating MD and 95% CI for A) Fractional area change, p-0.19; cell-based n=62; control n=62. B) End diastolic volume, p=0.52; cell-based n=110; control n=110. C) End systolic volume, p=0.96; cell-based n=110; control n=110.



Supplemental 10. A) Cell-based effect on human FAC.

		Experi	mental		(	Control				
Study	Total	Mean	SD	Total	Mean	SD	Mean Difference	MD	95%-CI	Weight
Ishigami 2014	7	119.70	94.20	7	113.20	106.10	i	6.50 [-	-98.61; 111.61]	6.9%
Ishigami 2014	7	99.80	65.80	7	104.60	76.20	<del></del>	-	-79.38; 69.78]	13.8%
Ishigami 2014	7	100.30	55.60	7	94.90	55.30		5.40	-52.69; 63.49]	22.7%
Ishigami 2017	17	115.80	170.70	17	113.00	113.40		2.80 [-	-94.62; 100.22]	8.1%
Ishigami 2017	17	100.40	192.50	17	99.50	107.20		0.90 [-	103.84; 105.64]	7.0%
Ishigami 2017	17	98.40	169.90	17	95.30	77.10		3.10 [	<b>-85.59</b> ; <b>91.79</b> ]	9.8%
Ishigami 2017	17	120.90	156.70	17	122.80	149.30	<del></del>	-1.90 [ <del>-</del>	104.79; 100.99	7.2%
Tarui 2015	7	160.40	128.60	7	139.00	114.80		– 21.40 [–	106.30; 149.10]	4.7%
Tarui 2015	7	150.20	110.30	7	112.20	83.10		38.00 [-	-64.30; 140.30]	7.3%
Tarui 2015	7	131.10	69.80	7	101.10	79.90		30.00 [-	-48.59; 108.59]	12.4%
Random effects mode	el 110			110				8.99 [	<b>-18.71</b> ; <b>36.69</b> ]	100.0%
Heterogeneity: $I^2 = 0\%$ , $\tau^2$	$^{2}$ = 0, $p$ =	: 1.00								
							-100 -50 0 50 100			

Supplemental 10. B) Cell-based effect on human EDV.

	Expe	rimental		Control				
Study	Total Mear	SD	Total Mean	SD	Mean Difference	MD	95%-CI Weig	jht
Ishigami 2014	7 60.30	41.30	7 56.90	80.40		3.40	[ -63.56; 70.36] 8.3	3%
Ishigami 2014	7 47.70	35.50	7 41.30	49.20	<del></del>	6.40	[ -38.54; 51.34] 18.4	1%
Ishigami 2014	7 48.50	42.30	7 38.00	33.30		10.50	[ -29.38; 50.38] 23.4	1%
Ishigami 2017	17 82.00	141.00	17 80.90	122.50		1.10	[-87.69; 89.89] 4.7	7%
Ishigami 2017	17 76.90	140.20	17 66.40	82.50		10.50	[ -66.83; 87.83] 6.2	<u>2</u> %
Ishigami 2017	17 55.90	138.10	17 50.30	80.40		5.60	[ -70.36; 81.56] 6.4	1%
Ishigami 2017	17 51.00	92.40	17 44.00	49.90		7.00	[-42.92; 56.92] 14.9	<del>)</del> %
Tarui 2015	7 27.40	72.50	7 58.90	49.70		-31.50	[ -96.62; 33.62] 8.8	3%
Tarui 2015	7 34.30	90.70	7 91.60	99.20		-57.30	[-156.87; 42.27] 3.7	7%
Tarui 2015	7 36.50	96.60	7 67.90	62.40		-31.40	[–116.59; 53.79] 5.1	1%
Random effects mode Heterogeneity: $I^2 = 0\%$ , $\tau^2$			110			-0.49	[ -19.78; 18.79] 100.0	)%
,	•			_	150–100 –50 0 50 100 15	0		

Supplemental 10. C) Cell-based effect on human ESV.

# 448 Supplemental 11. SYRCLE risk of bias for animal studies.

Author (Year)	Random sequence generation?	Groups similar at baseline?	Allocation concealed?	Animals randomly housed?	Blinding of caregivers and/or examiners?	Random selection for outcome assessment?	Blinding of outcome assessor?	Incomplete outcome data addressed?	Free from selective outcome reporting?	Free from other bias?
Agarwal (2016)	Yes	Yes	Yes	Unclear	Unclear	No	Yes	Yes	Yes	Yes
Albertario (2019)	Yes	Yes	No	Yes	Unclear	Unclear	Yes	Yes	Yes	Yes
Borenstein (2005)	Yes	Yes	Unclear	Unclear	Unclear	Unclear	Unclear	Yes	Yes	Yes
Brizard (2015)	Yes	Yes	Yes	Yes	Unclear	Unclear	Yes	Yes	Yes	Yes
Cao (2015)	Unclear	Yes	Unclear	Yes	Unclear	Unclear	Yes	Yes	Yes	Yes
Chery (2019)	Unclear	Yes	No	Unclear	Unclear	Unclear	Yes	Yes	Yes	Yes
Davies (2010)	Unclear	Yes	Unclear	Unclear	Unclear	Unclear	Unclear	Yes	Yes	No
Henning (2010)	Yes	Yes	Unclear	Unclear	Unclear	Unclear	Yes	Yes	Yes	Yes
Lambert (2015)	Unclear	Yes	Unclear	Unclear	Unclear	Unclear	Yes	Yes	Yes	Yes
Liu (2011)	Unclear	Yes	Unclear	Yes	Unclear	Unclear	Yes	Yes	Yes	Yes
Nana-Leventaki (2019)	Yes	Yes	Unclear	Yes	Unclear	Unclear	Yes	Yes	Yes	Yes
Schmuck (2019)	Yes	Yes	Unclear	Unclear	Unclear	Unclear	Yes	Yes	Yes	Yes
Sugiura (2016)	Yes	Yes	Unclear	Yes	Unclear	Unclear	Unclear	Yes	Yes	Yes
Trac (2018)	Yes	Yes	Unclear	Yes	Unclear	Unclear	Yes	Yes	Yes	Yes
Umar (2009)	Unclear	Yes	Unclear	Unclear	Unclear	Unclear	Yes	Yes	Yes	Yes
Wehman (2016)	Unclear	Yes	Unclear	Unclear	Unclear	Unclear	Yes	Yes	Yes	Yes
Wehman (2017)	Unclear	Yes	Unclear	Unclear	Unclear	Unclear	Yes	Yes	Yes	Yes
Yerebakan (2009)	Yes	Yes	Unclear	Unclear	Unclear	Unclear	Yes	Yes	Yes	Yes

#### Supplemental 12. ROBINS-I risk of bias for human studies.

457		
458		

456



Domains:

D1: Bias due to randomisation.

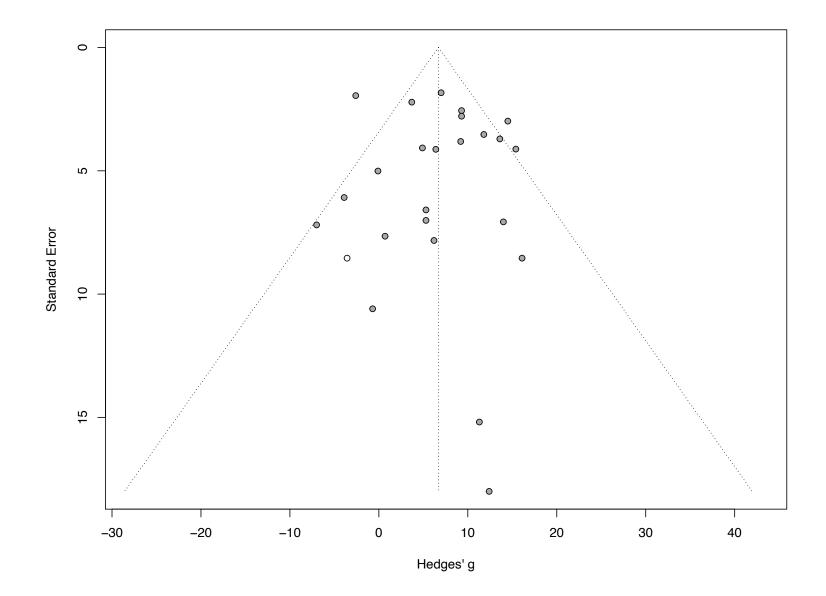
D2: Bias due to deviations from intended intervention.

D3: Bias due to missing data.

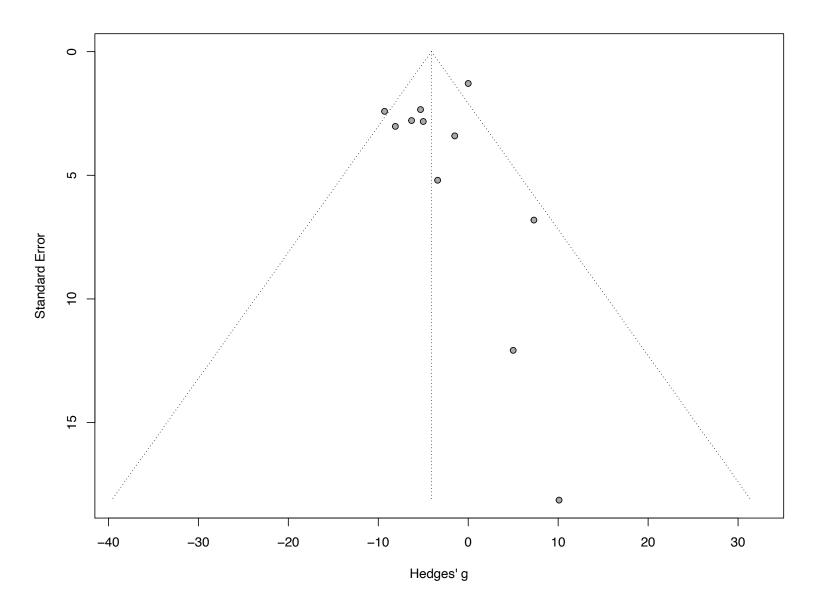
D4: Bias due to outcome measurement. D5: Bias due to selection of reported result. Judgement High

- Some concerns

+ Low



#### Supplemental 14. Funnel plot diagram for animal fractional shortening.



# 463 Supplemental Table 1. Animal study intervention characteristics.

Animal Characteristics	n (%)
Disease model	
RVHF	307 (80%)
DCM	45 (12%)
Autoimmune myocarditis	22 (6%)
Infant cardiopulmonary bypass	12 (3%)
Species	
Rats	120 (31%)
Sheep	118 (31%)
Rabbit	50 (13%)
Hamsters	45 (12%)
Swine	42 (11%)
Ram	11 (3%)
Age	
≤ 1 month	143 (37%)
>1 month	128 (33%)
Did Not Report	115 (30%)

Intervention Characteristics	n (%)
Cell route	
Intravenous	140 (36%)
Intramyocardial	135 (35%)
Graft/Patch/Sheet	61 (16%)
Intracoronary	34 (9%)
Epicardial	16 (4%)
Cell dose	
< 1M/kg	109 (28%)
≥ 1M and < 10M/kg	157 (41%)
≥ 10M/kg	120 (31%)
Cell source	
Cardiac	107 (28%)
Umbilical cord blood	93 (24%)
Bone marrow	80 (21%)
Adipose	70 (18%)
Thymus	25 (6%)
Skeletal muscle	11 (3%)
Timing	
Intraoperative and < 24 hours	79 (20%)
≥ 24 hours and < 1 week	0 (0%)
≥ 1 week and < 1 month	100 (26%)
≥ 1 month	177 (46%)
Did Not Report	30 (8%)

# **Supplemental Table 2. Adverse events by systems.**

Acute Adverse Events	
Systemic	
Allergic reaction / Anaphylaxis	Delayed Adverse Events
Death	Systemic
Elevated CRP	Late death
Fever	Cardiac
Hemodynamic instability / Hypotension	Late heart failure
Malignancy	
Tumor formation	Unplanned Interventions
Respiratory	BCPS or TCPC take down
Bronchitis	Catheterizations
Chest tube	Heart transplant
Death - Respiratory	Intubation
Pleural effusion	Pacemaker implantation
Pneumonia	·
Cardiac	Unplanned Hospitalizations
APCA coil occlusion	CCU admission
Arrythmia	ED visit
Bradycardia	General wards/floor admission
Cardiac tamponade	Rehospitalization for heart failure
Cardiac-related pneumothorax	
Cardiopulmonary resuscitation	
Coronary spasm	APCA = aortopulmonary collateral artery
Death - Cardiac	BCPS = bidirectional cavopulmonary shunt
ECMO	CCU = cardiac/coronary care unit
Epicardial bleed	ED = emergency department
Heart failure	TCPC = total cavopulmonary connection
Myocardial ischemia	
Palpitations	
Valve malfunction	
Neurological	
Seizure	
Stroke	
Hematological	
Embolism	
Thromboembolic events	
GI	4
Cirrhosis	4
Protein-losing enteropathy	-
Renal Danel deterioration	-
Renal deterioration	_
Infectious disease	$\dashv$
Infection	

# **Supplemental Table 3. Animal adverse events.**

Adverse Event	Cor	ntrol	Cell-k	Overall	
Adverse Event	Events	Total	Events	Total	Peto OR (95% CI)
Cardiac	13	73	8	74	0.48 (0.17, 1.33)
Respiratory	5	69	8	67	2.29 (0.67, 7.84)
GI	0	69	0	67	-
Hematologic	0	69	0	67	-
Infectious disease	0	69	0	67	-
Systemic	1	69	0	67	0.44 (0.02, 12.01)
Overall	19	418	16	409	0.89 (0.43, 1.83)

# Supplemental Table 4. Human study intervention characteristics.

Human Characteristics	n (%)		
Disease			
SV	142 (66%)		
HLHS	41 (19%)		
DCM	32 (15%)		
Cell route			
Intracoronary	204 (95%)		
Intramyocardial	11 (5%)		
Cell dose			
< 1M/kg	170 (79%)		
≥ 1M and < 10M/kg	35 (16%)		
≥ 10M/kg	0 (0%)		
Did Not Report	10 (5%)		
Cell source			
Cardiac	170 (79%)		
Bone marrow	34 (16%)		
Umbilical cord blood	11 (5%)		

# 471 Supplemental Table 5. Human adverse events.

	Control		Cell-based		Overall	
Adverse Event	Events	Total	Events	Total	Peto OR (95% CI)	
Cardiac	81	101	41	82	0.11 (0.05, 0.23)	
Respiratory	5	101	0	82	0.16 (0.03, 0.95)	
GI	2	101	0	82	0.14 (0.01, 2.16)	
Hematologic	7	101	2	82	0.44 (0.11, 1.75)	
Infectious disease	1	101	0	82	0.19 (0.00, 10.05)	
Systemic	2	101	1	82	0.74 (0.07, 7.54)	
Overall	98	606	44	492	0.17 (0.09, 0.30)	

# Supplemental Table 6. Clinical trials, ongoing.

Sponsor	Disease	Methods	Participants	Interventions	Comparison	Outcome	Notes
NCT03779711	HLHS	Phase II treatment given at time of stage II surgical repair	Children <9 months of age with HLHS or HLHS variant with single ventricular dependent CHD having undergone Stage I surgical repair and Stage II surgical repair	Biologic: Autologous UCB derived mononuclear cells Procedure: Stage II surgical repair	Stage II surgical repair alone	Efficacy	Recruiting
NCT03525418	HLHS	Phase I/II: First treatment given at time of stage II surgical repair  Second treatment given at time of stage II surgical repair vs. placebo	HLHS (all types) requiring stage II surgical intervention	Biologic: Bone marrow derived MSCs Procedure: Stage II surgical repair	Stage II surgical repair alone	Safety and Efficacy	Recruiting
NCT03431480	HLHS	Phase I open label safety study	Male and females with antenatally diagnosed HLHS (all types requiring Norwood operation)	Biologic: Autologous human placental cord blood mononuclear cells Procedure: Stage I surgical repair	N/A	Safety and Efficacy	Recruiting
NCT03079401	AV Canal Defects	Phase I/II	Patient with a history of single ventricle palliation undergoing bidirectional Glenn with LV recruitment procedures or those patients undergoing LV recruitment procedures	Biologic: Mesenchymal progenitor cells	Surgical repair alone	Safety and Efficacy	Recruiting
NCT02781922	HLHS SV	Phase III single blind parallel group study	Functional single ventricle patient with HF scheduled for stage 2 (Glenn) or stage 3 (Fontan) surgery	Biologic: Autologous cardiac stem cells	Surgical repair alone	Safety and Efficacy	Unknown

			EF < 55%				
NCT01883076	HLHS	Phase I	Individuals with HLHS who have undergone Stage I surgical palliation and undergoing planned Stage II Palliative Glenn Surgery	Biologic: autologous umbilical cord blood cells	N/A	Safety	Recruiting